<110> Rigshospitalet

SEQUENCE LISTING

Henrik Leffers Anne Mette Buhl Hertz Jørgen Kjems <120> Methods and kits for diagnosing and treating B-cell Chronic lymphocytic leukemia (B-CLL) <130> P34546PC01 <150> DK/PA 200201792 <151> 2002-11-19 <160> 43 <170> FastSEQ for Windows Version 4.0 <210> 1 <211> 19999 <212> DNA <213> Homo Sapiens <220> <221> gene <222> 40000-60000 <223> Sequence of ac063949.emhum <400> 1 cacacgtagg ctacgagtgg ccctcagcct gcctcatcat ggacctgtgt tataataaat 60 atgtttaatt gtgctgtttt cttatagagg aaagtcctga tgttagttgc cttgaagtca 120 gacacccaga gagaatcaca ggttttcaga ttaattcatc gcttgattct tatccctgaa 180 gtcatatctc tggatctctg gttctcacat tataaatttc aatgattctt tttctatatg 240 gccatgtcat tcatatcctg tgtaatatgg ggaaactgag gtatgaatga catcattcaa 300 aaagcacctg caatttttct ttgccaagca cttacagctt tttctcatgt tgctttcaaa 360 aagtcattga aatattgttc acatattttg cagatgagga aatgaatatt caaatgcatt 420 aggtatettg tecaagttet tacagecaga aagtagagaa atgaatttga attacaaate 480 ttctacctct tggcttatgc tcttttcatg acactgggaa taaatgtctg aacaagcatg 540 acttcatgtt tcaactattt atcaaatact tgttttctac taagatcttg cactcactca 600 gtgggatccc ctgaagcctg ctgattattt gtcctttggc atttatcact ctctgtggga 660 ccttactctc ctatggtaaa gttttattgt tattaaaagt attatttgac aataaatgta 720 gaaatcctac agatcatact caacaacatg tctaatgtca gcacacaatg tctaacaatc 780 atttatgaat actttatgtc aaacataagc aataacctaa ttaaggaagg tatttttaat 840 aaattgacac tttttgacat aaccatattt caagtggctc cattgttttg tttatttatt 900 tatttattta tttatttatt tatttttgag aaagggtctc actctgttgc ccagactgga 960 gtgcagtggc aacatcatag ctcactacag cctcgacctc tctgggctca agcaatcctc 1020 ccatctcagc ttcccaagta gctgggacta caggtgtgta ccatcatgcc aggctaattt 1080 ttcgtatttt gtagagacgg ggttttgcct ggtcgtccgg gttggtctca aactcctggg 1140 tgttccgccc accttggcct cccaaagtgc tgggattata ggcatgagcc tcaagtggct 1200 actttttagg gttgaaattt atattgactg tcaactagct tccctagtta gtatttggga 1260 tctgctaact aatttatatt accatccaac ttgtcaacat ttgttgaaat ataactgtcc 1320 tcactttttt tgtgtgaaca ttgaatacac tttcagacta aatttggttt attacttaat 1380 gtcttattct ttattagagt taataatatt tcttaatact ttgccttcca caaatgaata 1440 acttgtttgt gatggctacc tcttttttc tcttagcctg tcacaggtat tatggataaa 1500 aattagcacg gctgggcaaa aacaatgaaa gaaatacact tgcctgggaa agctggggag 1560 gggtaaatga atataattca aaataccata tatttattca acactgttgg aatatatgtc 1620 ctgttggaaa tgtaaaagtg acatatgttc tcttcctggg tctcagactt ttaggatcta 1680 gttgagggaa ctggacttat acacaaaata caattcaaca acattatgag ctagaaaatc 1740

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WO 2004/046376 PCT/DK2003/000794

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<223> Corresponds to a cDNA detected by cDNA cloning. It
      includes three exons, the first and third are
      identical to the two exons in Seq_ID_06.doc. It is
      transcribed from the +strand of Seq ID 06.doc
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34/48

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      includes two exons, the last is also present as
      "exon-2" in a human cDNA clone (sequence acc. no.
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35/48

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<223> Corresponds to a human cDNA sequence (sequence
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<211> 3931

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<213> Homo sapiens

<220>

<221> mRNA

<222> (0)...(0)

<223> Corresponds to a cDNA detected partly by cDNA cloning, partly by PCR analysis. It includes two exons, the last is identical to the last exon in Seg ID 06 and Seg ID 07.doc. It is transcribed from the +strand of Seq_ID_06.doc

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<222> (0)...(0)
<223> Corresponds to a cDNA detected by cDNA cloning. It includes one exon. The sequence includes the last two exons in Seq_ID_07.doc. It is transcribed from the +strand of Seq_ID_06.doc

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47/48

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WO 2004/046376 PCT/DK2003/000794

48/48

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